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Amendments to the Claims:

This listing of claims will replace all prior versions, and listings, of claims in the application.

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Listing of Claims:

1-51 (Cancelled).

52 (Original). A method of producing in a eukaryotic cell at least one protein comprising at least one unnatural amino acid, the method comprising:

growing, in an appropriate medium, a eukaryotic cell that comprises a nucleic acid that comprises at least one selector codon and encodes the protein; wherein the medium comprises an unnatural amino acid and the eukaryotic cell comprises:

an orthogonal tRNA (O-tRNA) that functions in the cell and recognizes the selector codon; and,

an orthogonal aminoacyl tRNA synthetase (O-RS) that preferentially aminoacylates the O-tRNA with the unnatural amino acid, wherein the O-RS comprises an amino acid sequence that corresponds to SEQ ID NO.: 48 – 53.

53 (Currently Amended). A method of producing in a eukaryotic cell at least one protein comprising at least one unnatural amino acid and modifying said protein, the method comprising:

growing, in an appropriate medium, a eukaryotic cell that comprises a nucleic acid that comprises at least one selector codon and encodes the protein; wherein the medium comprises the unnatural amino acid and the eukaryotic cell comprises an orthogonal tRNA (O-tRNA) that functions in the cell and recognizes the selector codon and an orthogonal aminoacyl tRNA synthetase (O-RS) that preferentially aminoacylates the O-tRNA with the unnatural amino acid, wherein the O-RS is selected from:

- (i) a p-propargyloxyphenylalanine O-RS that preferentially aminoacylates the O-tRNA with p-propargyloxyphenylalanine, and
- (ii) a p-azido-L-phenylalanine O-RS that preferentially aminoacylates the O-tRNA with p-azido-L-phenylalanine, the p-azido-L-phenylalanine O-RS comprising:

(i) an amino acid sequence set forth in any one of SEQ ID NOs: 48 – 53[[,]] and conservative variants thereof, which wherein the conservative variants are at least 90% 98% identical to SEQ ID NOs 48 – 53, or

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> (ii) an amino acid sequence that is a conservative variant of SEQ ID NO: 2, which conservative variant is at least 98% identical to SEQ ID NO: 2 that of a naturally occurring tyrosyl aminoacyl tRNA synthetase (TyrRS) and comprises two or more amino acids selected from the group consisting of: glycine, serine, or alanine at a position corresponding to Tyr37 of E. coli TyrRS; aspartate at a position corresponding to Asn126 -of-E. coli TyrRS; asparagine at a position corresponding to Asp182 of E. coli TyrRS; alanine, or valine, at a position corresponding to Phe183 of E. coli TyrRS; and, methionine, valine, cysteine, or threonine, at a position corresponding to Leu186 of E. coli TyrRS:

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incorporating into the protein the unnatural amino acid in the eukaryotic cell, wherein the unnatural amino acid comprises a first reactive group; and.

contacting the protein with a molecule that comprises a second reactive group; wherein the first reactive group reacts with the second reactive group to attach the molecule to the unnatural amino acid through a [3+2] cycloaddition, thereby modifying the protein.

- 54 (Original). The method of claim 53, wherein the molecule is a dye, a polymer, a derivative of polyethylene glycol, a photocrosslinker, a cytotoxic compound, an affinity label, a derivative of biotin, a resin, a second protein or polypeptide, a metal chelator, a cofactor, a fatty acid, a carbohydrate, or a polynucleotide.
- 55 (Cancelled).
- 56 (Previously Presented). The method of claim 53, wherein the O-RS is the ppropargyloxyphenylalanine O-RS, the unnatural amino acid is p-propargyloxyphenylalanine, the first reactive group is an alkynyl moiety and the second reactive group is an azido moiety.
- 57 (Previously Presented). The method of claim 53, wherein the unnatural amino acid comprises a p-propargyloxyphenylalanine.
- 58 (Currently Amended). The method of claim 53, wherein the O-RS is the p-azido-Lphenylalanine O-RS, the unnatural amino acid is p-azido-L-phenylalanine, the first reactive group is an azido moiety, and the second reactive group is an alkynyl moiety.
- 59 (Previously Presented). The method of claim 53, wherein the unnatural amino acid comprises a p-azido-L-phenylalanine.
- 60 61 (Cancelled).

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- 62 (Withdrawn). The method of claim 53, wherein the O-RS is the *p*-propargyloxyphenylalanine O-RS which *p*-propargyloxyphenylalanine O-RS comprises an amino acid sequence set forth in any one of SEQ ID NOS: 54-63, and conservative variants thereof, which conservative variants are at least 90% identical to that of a naturally occurring tyrosyl aminoacyl-tRNA synthetase (TyrRS) and comprise two or more amino acids selected from the group consisting of: glycine, serine, or alanine at a position corresponding to Tyr37 of *E. coli* TyrRS; aspartate at a position corresponding to Asn126 of E. coli TyrRS; asparagine at a position corresponding to Asp182 of *E. coli* TyrRS; alanine, or valine, at a position corresponding to Phe183 of *E. coli* TyrRS; and, methionine, valine, cysteine, or threonine, at a position corresponding to Leu186 of *E. coli* TyrRS.
- 63 (Currently Amended). A method of producing in a yeast cell at least one protein of interest comprising at least one unnatural amino acid selected from p-azido-L-phenylalanine and p-propargyloxyphenylalanine, the method comprising growing, in an appropriate medium, the yeast cell, wherein the medium comprises the unnatural amino acid, and wherein the yeast cell comprises:
- (a) a nucleic acid encoding said protein of interest and comprising at least one selector codon; and;
- (b) an orthogonal tRNA (O-tRNA) that functions in the cell and recognizes the selector codon, wherein the O-tRNA is derived from an *Escherichia coli* tRNA; and,
- (c) an orthogonal aminoacyl tRNA synthetase (O-RS) that preferentially aminoacylates the O-tRNA with the unnatural amino acid, wherein the O-RS is derived from an *Escherichia coli* aminoacyl tRNA synthetase.
- 64 (Currently Amended). A method of producing in a eukaryotic cell at least one protein comprising at least one unnatural amino acid selected from p-azido-L-phenylalanine and p-propargyloxyphenylalanine, the method comprising:
- (a) growing, in an appropriate medium, a eukaryotic cell that comprises a nucleic acid that comprises at least one selector codon and encodes the protein; wherein the medium comprises the unnatural amino acid and wherein the eukaryotic cell comprises an orthogonal tRNA (O-tRNA) that functions in the cell and recognizes the selector codon and an orthogonal aminoacyl tRNA synthetase (O-RS) that preferentially aminoacylates the O-tRNA with the unnatural amino acid, wherein the O-RS is selected from:
 - (i) a p-propargyloxyphenylalanine O-RS that preferentially aminoacylates the OtRNA with p-propargyloxyphenylalanine, or

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- (ii) a p-azido-L-phenylalanine O-RS that preferentially aminoacylates the O-tRNA with p-azido-L-phenylalanine, the p-azido-L-phenylalanine O-RS comprising:
 - (i) an amino acid sequence set forth in any one of SEQ ID NOs: 48 53[[,]] and conservative variants thereof, which wherein the conservative variants are at least 90% 98% identical to SEQ ID NOs 48 53, or
 - (ii) an amino acid sequence that is a conservative variant of SEQ ID NO: 2, which conservative variant is at least 98% identical to SEQ ID NO: 2 that of a naturally occurring tyrosyl aminoacyl tRNA synthetase (TyrRS) and comprises two or more amino acids selected from the group consisting of: glycine, serine, or alanine at a position corresponding to Tyr37 of E. coli TyrRS; asparate at a position corresponding to Asn126 of E. coli TyrRS; asparagine at a position corresponding to Asp182 of E. coli TyrRS; alanine, or valine, at a position corresponding to Phe183 of E. coli TyrRS; and, methionine, valine, cysteine, or threonine, at a position corresponding to Leu186 of E. coli TyrRS; and,
 - (b) incorporating into the protein the unnatural amino acid in the eukaryotic cell.